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P	⊠		US 6383754 B1	07	43	Binary encoded sequence , tags	435/6	435/91.1; 435/91.2; 436/94	
2			US 6380369 B1	20020430	68	Human DNA mismatch repair proteins	536/23.1	435/6	
ω	Ø		US 6361947 B1	20020326	34	Complexity management and analysis of genomic 4	435/6	536/23.1	
4	×		US 6309823 B1	20011030	90	Arrays of nucleic acid probes for analyzing biotransformation genes and methods of using the same	435/6	422/50; 422/68.1	
vi	×		US 6261782 B1	20010717	61	Fixed address analysis of sequence tags	435/6	435/91.2; 536/23.1; 536/24.2; 536/24.3	
9	⊠		US 6239264 B1	20010529	637	Genomic DNA sequences of ashbya gossypii and uses thereof	536/23.1	435/320.1; 536/24.3; 536/24.32	
7	Ø		US 6221585 B1	20010424	54	Method for identifying genes underlying defined phenotypes	435/6	435/91.2	

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н	Kaufman, Joseph C. et al.								US 6383754	
2	Adams, Mark D. et al.	⊠							US 6380369	
ω	Dong, Shoulian et al.								US 6361947	
4	Cronin, Maureen T. et al.								US 6309823	
U	Lizardi, Paul M. et al.								US 6261782	
D	Philippsen, Peter et al.								US 6239264	
7	Iris, Francois J. -M. et al.								US 6221585	

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- AN 1999:57842 BIOSIS
- DN PREV199900057842
- TI The role of mismatch repair in the prevention of base pair mutations in Saccharomyces cerevisiae.
- AU Earley, Marie C.; Crouse, Gray F. (1)
- CS (1) Dep. Biol., Emory Univ., Atlanta, GA 30322 USA
- Proceedings of the National Academy of Sciences of the United States of America, (Dec. 22, 1998) Vol. 95, No. 26, pp. 15487-15491.
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- AN 1997:296017 BIOSIS
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- AU Petes, Thomas D. (1); Greenwell, Patricia W.; Dominska, Margaret
- CS (1) Dep. Biol., Univ. North Carolina, Chapel Hill, NC 27599-3280 USA
- SO Genetics, (1997) Vol. 146, No. 2, pp. 491-498. ISSN: 0016-6731.
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- AU Greene, Christopher N.; Jinks-Robertson, Sue (1)
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- SO Molecular and Cellular Biology, (1997) Vol. 17, No. 5, pp. 2844-2850. ISSN: 0270-7306.
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- L5 ANSWER 5 OF 6 BIOSIS COPYRIGHT 2002 BIOLOGICAL ABSTRACTS INC.
- AN 1995:157267 BIOSIS
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- TI The Saccharomyces cerevisiae Msh2 protein specifically binds to duplex oligonucleotides containing mismatched DNA base pairs and insertions.
- AU Alani, Eric; Chi, Nai-Wen; Kolodner, Richard (1)
- CS (1) Div. Cellular Molecular Biology, Dana Farber Cancer Inst., Dep.
  Biological Chemistry Molecular Pharmacology, Harvard Med. Sch., Boston, MA
  02115 USA
- SO Genes & Development, (1995) Vol. 9, No. 2, pp. 234-247. ISSN: 0890-9369.
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- L5 ANSWER 6 OF 6 CAPLUS COPYRIGHT 2002 ACS
- AN 2001:636237 CAPLUS
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- TI Methods and uses thereof for generating hypermutable yeast for mutagenesis

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    Kinzler, Kenneth W.
    The Johns Hopkins University, USA
PΑ
SO
    PCT Int. Appl., 59 pp.
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LA
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    PATENT NO.
                    KIND DATE
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            SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN,
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L1 417 S MSH3
L2 64 S L1 AND PLANT
L3 61 DUP REM L2 (3 DUPLICATES REMOVED)
L4 58 S L3 AND MISMATCH
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6 S L4 AND (MISMATCH REPAIR SYSTEM)

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